

Survey: **KU3/25**
month: **July 2025**
Participant-No: **9900540**
valid from: **05.07.25**



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Všeobecná fakultní nemocnice v Praze
Centrální Laborator na ULBLD
Prof. MUDr. Zima, RNDr. Benáková
U Nemocnice 499/2, budova A 7, 2. patro
12808 PRAHA 2
CZECH REPUBLIC



Bonn, 25. July 2025

Certificate

We confirm that you have participated in the survey for clinical chemical analytes in urine.
This certificate is according to the current version of the RiliBÄK valid until the end of January 2026.
You have met the requirements of the survey for the following analytes:

glucose	(1)	uric acid	(1)	urea	(3)
creatinine	(1)	total protein	(5)	a1-microglobulin	(2)
albumin	(2)	IgG	(1)	calcium	(6)
chloride	(4)	potassium	(2)	magnesium	(1)
sodium	(2)	phosphorus, inorg.	(2)	osmolality	(1)
proteinuria	(1)	pH ST	(2)	protein ST	(2)
glucose ST	(2)	hemoglobin ST	(2)	leukocytes ST	(2)
keton ST	(2)	nitrite ST	(2)	bilirubin ST	(2)
specific grav. ST	(2)	urobilinog. ST	(2)		

Prof. Dr. Dr. K. P. Kohse
EQA scheme director

Dr. Anja Kessler
Head of Reference Institute

Dr. Marika Enders
EQAS-Board

The number in parentheses characterizes the analytical method used.
The assignment of the number to the respective method and/or the respective instrument
is to be taken from the total evaluation.

This certificate is valid in conjunction with the final report dated 15.07.25.
This is available for download in the RfB-Online system.

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Centralni Laborator na ULBLD
Prof. MUDr. Zima, RNDr. Benáková
U Nemocnice 499/2, budova A 7, 2. patro
12808 PRAHA 2
CZECH REPUBLIC



Bonn, 25. July 2025

Participation certificate

We confirm that you have participated in the survey for clinical chemical analytes in urine.

The analytes determined by you are as follows:

glucose	(1)	uric acid	(1)	urea	(3)
creatinine	(1)	citrate	(1)	oxalate	(1)
total protein	(5)	a1-microglobulin	(2)	albumin	(2)
IgG	(1)	calcium	(6)	chloride	(4)
potassium	(2)	magnesium	(1)	sodium	(2)
phosphorus, inorg.	(2)	osmolality	(1)	proteinuria	(1)
pH ST	(2)	protein ST	(2)	glucose ST	(2)
hemoglobin ST	(2)	leukocytes ST	(2)	keton ST	(2)
nitrite ST	(2)	bilirubin ST	(2)	specific grav. ST	(2)
urobilinog. ST	(2)	diagn. graph A	(1)	diagn. graph B	(1)

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Listing and Evaluation of all your results

Explanations

Certificate

A certificate is issued (given) for an analyte only if the basis for an evaluation of the accuracy is given by the guidelines of the German Medical Association and/or if an evaluation is possible in analogy to these guidelines (see comments on the evaluation) and, if both results for an analyte are within the given acceptance limits. (marked as '+' below C)

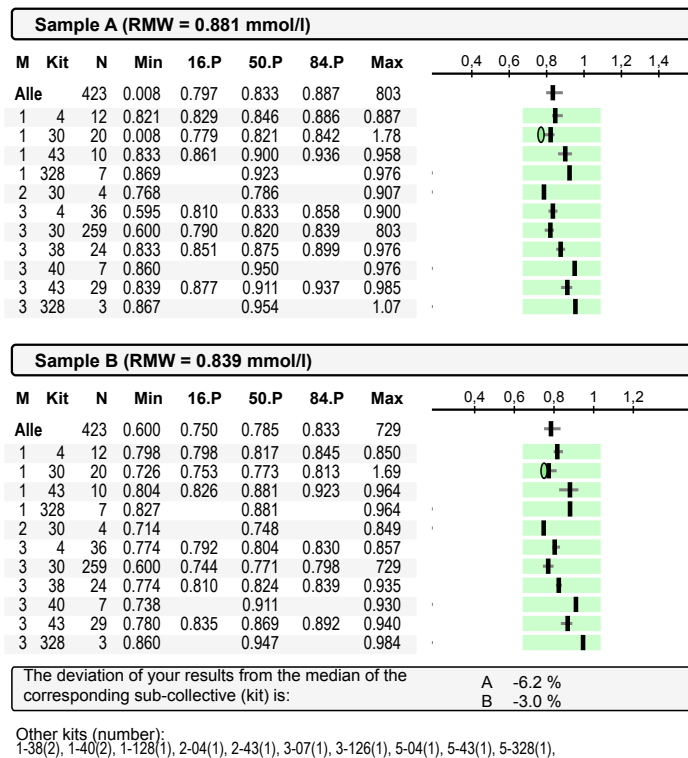
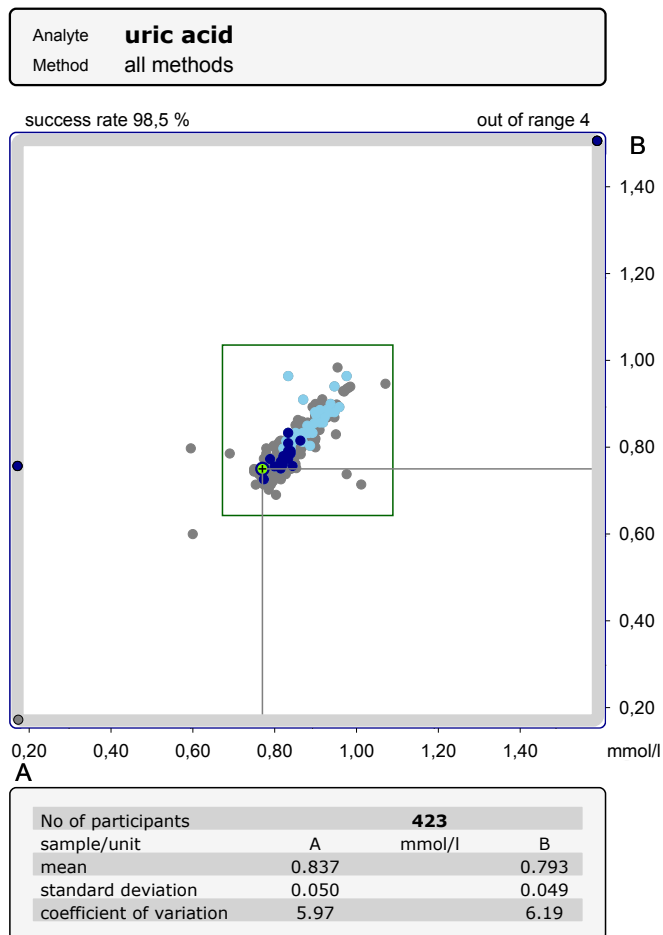
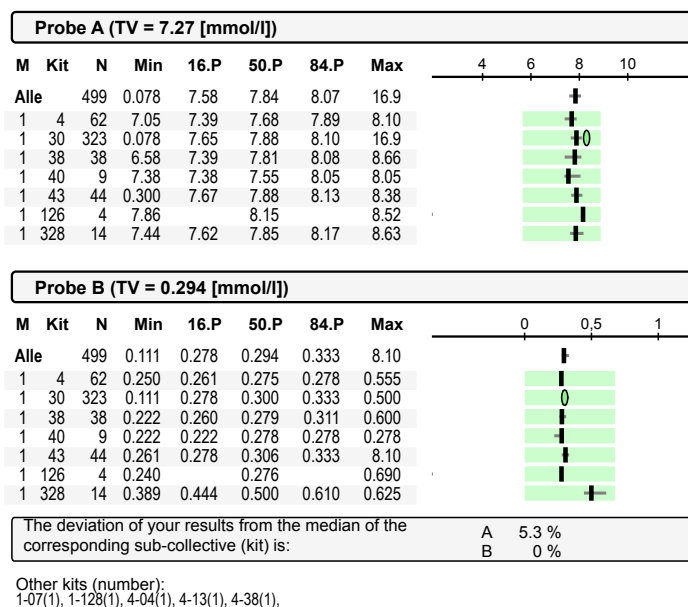
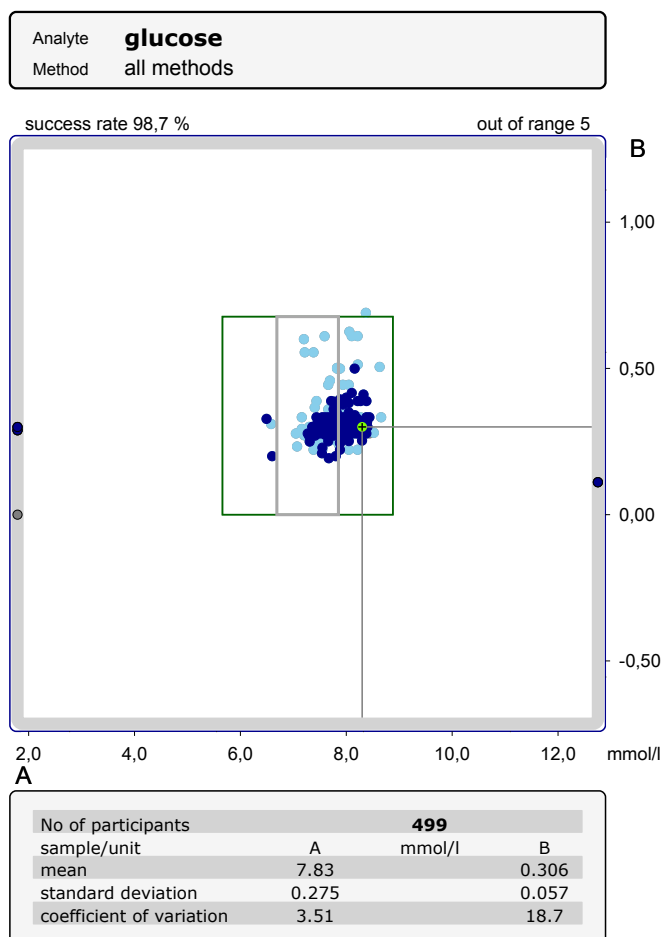
Certificate of participation

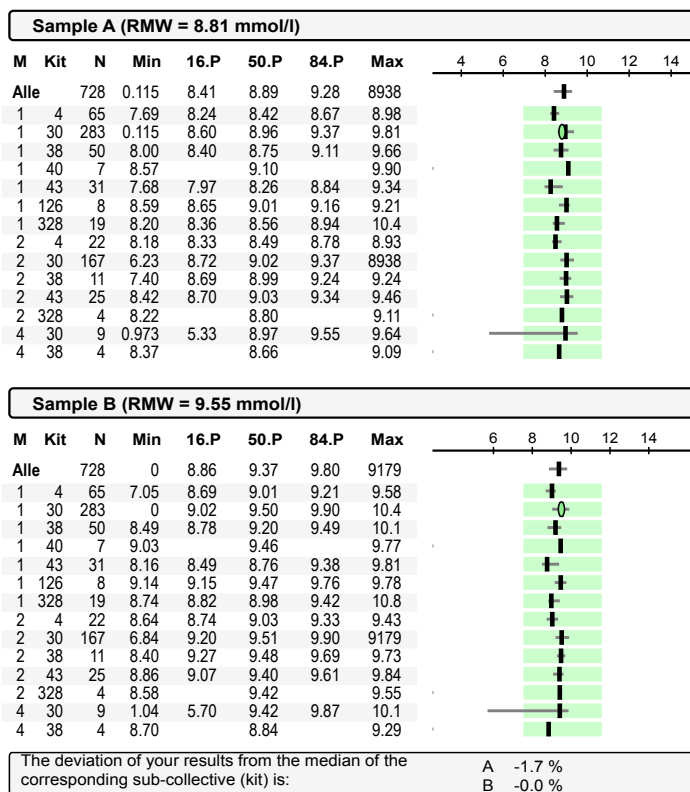
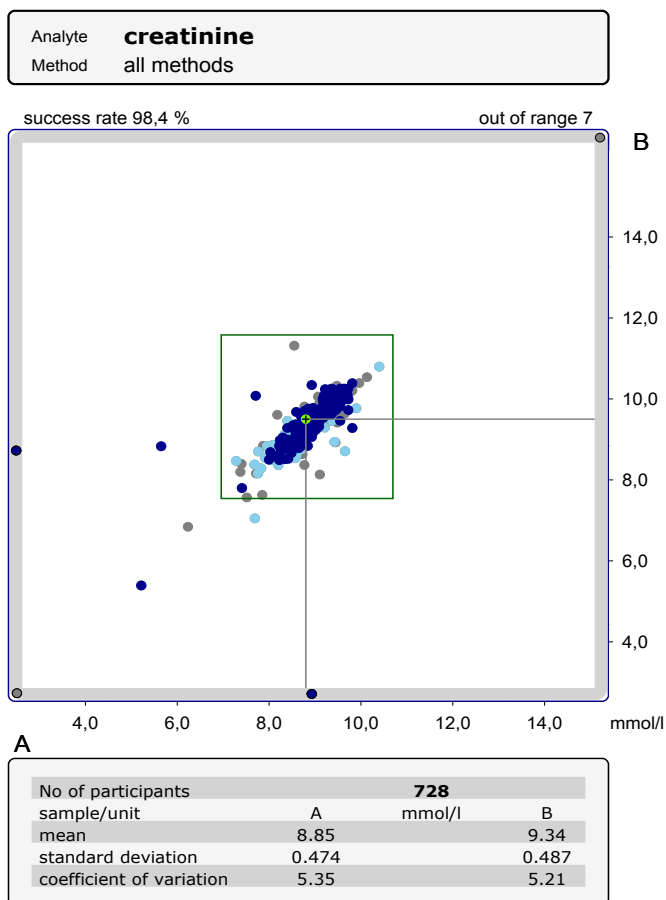
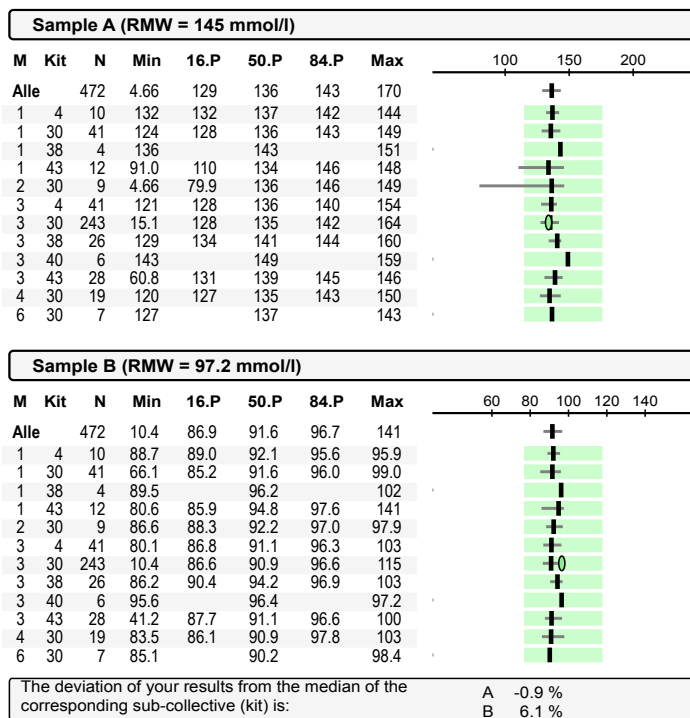
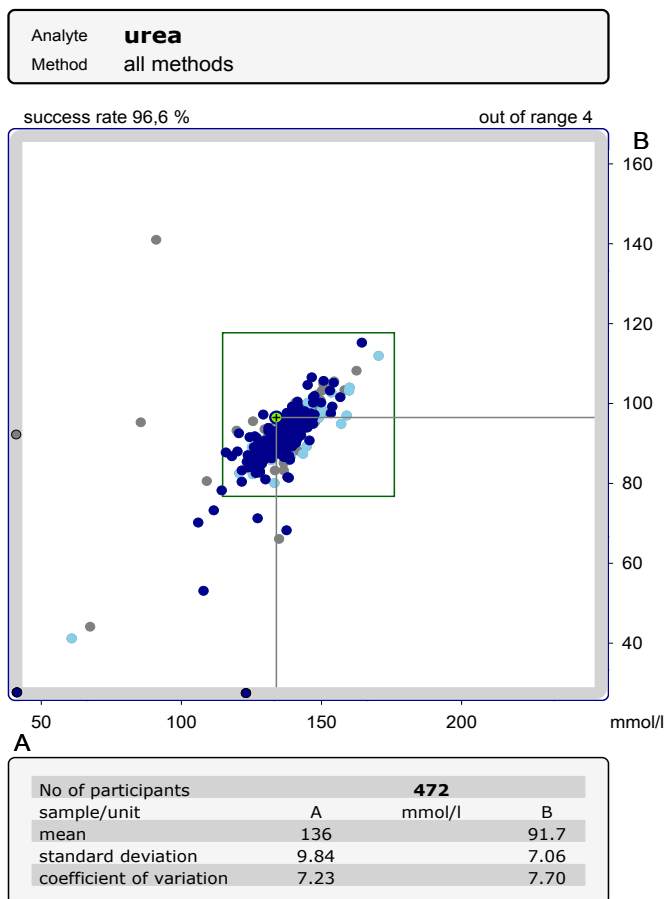
In the participation certificate all analytes which are included in the list on this page are listed.

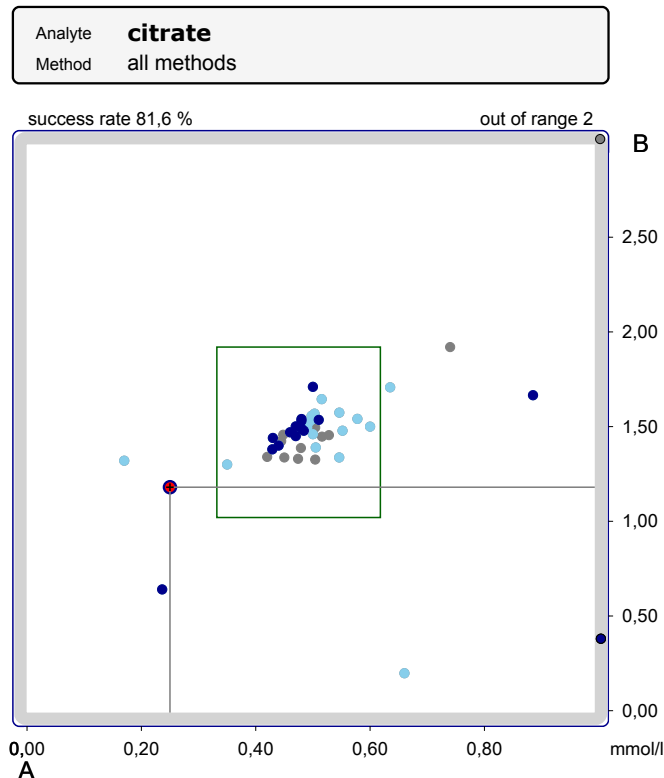
If all analytes are listed on the certificate no participation certificate is printed.

Legend:	C=Certification, M=No of method, R=your result, D=difference (R-T) Dmax= maximum allowable amount of difference in measurement, partly given by annex 1 of the guideline of the BÄK. T = target value, either reference method value or assigned value, LL UL = lower resp. upper limit	Certification: + = fulfilled (quotient D/Dmax <= 1.0) - = not fulfilled (quotient D/Dmax > 1.0 ± = certification cancelled because of technical and/or analytical reasons
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	C	M	R	D/Dmax	T	LL	UL		C	M	R	D/Dmax	T	LL	UL
glucose	+	1	A 8.30	0.64	7.27	5.66	8.88	glucose ST	+	2	A Ch. 3		Ch. 3	Ch. 2	Ch. 3
[mmol/l]			B 0.300	0.01	0.294	0	0.678	[ohne]			B Ch. 1		Ch. 1	Ch. 1	Ch. 1
uric acid	+	1	A 0.770	-0.53	0.881	0.672	1.09	hemoglobin ST	+	2	A Ch. 1		Ch. 1	Ch. 1	Ch. 1
[mmol/l]			B 0.750	-0.45	0.839	0.642	1.04	[ohne]			B Ch. 1		Ch. 1	Ch. 1	Ch. 1
urea	+	3	A 134	-0.35	145	114	176	leukocytes ST	+	2	A Ch. 1		Ch. 1	Ch. 1	Ch. 1
[mmol/l]			B 96.5	-0.03	97.2	76.7	118	[ohne]			B Ch. 1		Ch. 1	Ch. 1	Ch. 1
creatinine	+	1	A 8.80	-0.01	8.81	6.95	10.7	keton ST	+	2	A Ch. 1		Ch. 1	Ch. 1	Ch. 1
[mmol/l]			B 9.50	-0.02	9.55	7.54	11.6	[ohne]			B Ch. 1		Ch. 1	Ch. 1	Ch. 1
citrate	-	1	A 0.250	-1.57	0.475	0.332	0.618	nitrite ST	+	2	A neg		neg	neg	neg
[mmol/l]			B 1.18	-0.64	1.47	1.02	1.92	[ohne]			B neg		neg	neg	neg
oxalate	-	1	A 190	0.72	143	78.6	208	bilirubin ST	+	2	A Ch. 1		Ch. 1	Ch. 1	Ch. 1
[µmol/l]			B 187	1.45	113	62.1	164	[ohne]			B Ch. 1		Ch. 1	Ch. 1	Ch. 1
total protein	+	5	A 0.600	0.07	0.590	0.448	0.732	specific grav. ST	+	2	A 1.010	0	1.010	1.000	1.020
[g/l]			B 0.030	-0.17	0.036	0	0.100	[g/ml]			B 1.005	-0.50	1.010	1.000	1.020
a1-microglobulin	+	2	A 55.0	-0.10	57.5	32.2	82.8	urobilinog. ST	+	2	A Ch. 1		Ch. 1	Ch. 1	Ch. 1
[mg/l]			B 2.20	-0.56	5.00	0	20.0	[ohne]			B Ch. 1		Ch. 1	Ch. 1	Ch. 1
albumin	+	2	A 464	0.18	443	327	559	diagn. graph A	±	1	A 466				
[mg/l]			B 10.8	0.01	10.7	0	20.0	[mg/g Krea]			B 55.3				
IgG	+	1	A 44.4	-0.01	44.5	31.1	57.9	diagn. graph B	±	1	A 10.1				
[mg/l]			B 3.00	0	3.00	0	10.0	[mg/g Krea]			B 2.05				
calcium	+	6	A 0.980	0.27	0.935	0.776	1.10								
[mmol/l]			B 1.41	-0.04	1.42	1.17	1.67								
chloride	+	4	A 71.0	0	71.0	61.0	81.0								
[mmol/l]			B 46.1	-0.06	46.5	39.9	53.1								
potassium	+	2	A 24.0	0	24.0	20.4	27.6								
[mmol/l]			B 24.4	0.05	24.2	20.5	27.9								
magnesium	+	1	A 2.26	0.01	2.25	1.66	2.84								
[mmol/l]			B 1.75	0.13	1.69	1.25	2.13								
sodium	+	2	A 94.0	-0.09	95.0	83.6	107								
[mmol/l]			B 61.4	-0.21	63.0	55.4	70.6								
phosphorus, inorg.	+	2	A 9.40	-0.26	9.91	7.91	12.0								
[mmol/l]			B 7.20	-0.26	7.59	6.07	9.11								
osmolality	+	1	A 370	-0.13	376	330	422								
[mosm/kg]			B 257	-0.28	266	234	298								
proteinuria	+	1	A Ch. 13		Ch. 13,										
[ohne]			B Ch. 11		Ch. 11,										
pH ST	+	2	A 7		7,5	7	8								
[ohne]			B 6,5		7,5	6,5	8								
protein ST	+	2	A Ch. 4		Ch. 4	Ch. 3	Ch. 4								
[ohne]			B Ch. 1		Ch. 1	Ch. 1	Ch. 1								







No of participants	49		
sample/unit	A	mmol/l	B
mean	0.490		1.43
standard deviation	0.112		0.253
coefficient of variation	22.7		17.7

Sample A [mmol/l]

M	Kit	N	Min	16.P	50.P	84.P	Max
Alle	49	0.170	0.440	0.490	0.578	10.7	
1	30	7	0.490		0.503	0.635	
1	38	3	0.490		0.505	0.505	
1	99	16	0.236	0.379	0.475	0.615	1.08
1	148	6	0.350		0.531	0.578	
4	99	9	0.440	0.446	0.479	4.72	10.7

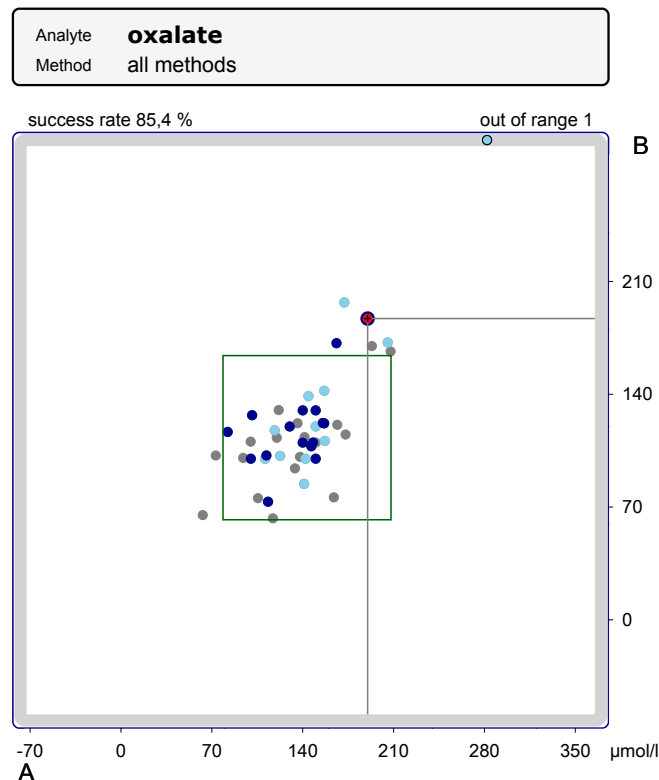
Sample B [mmol/l]

M	Kit	N	Min	16.P	50.P	84.P	Max
Alle	49	0.198	1.33	1.47	1.57	23.3	
1	30	7	1.34		1.50	1.71	
1	38	3	1.39		1.49	1.54	
1	99	16	0.380	1.03	1.47	1.58	1.71
1	148	6	1.30		1.55	1.64	
4	99	9	1.33	1.33	1.46	10.5	23.3

The deviation of your results from the median of the corresponding sub-collective (kit) is:

A-47.37 %
B-19.96 %

Other kits (number):
1-126(1), 1-168(1), 2-99(2), 3-99(1), 4-12(1), 4-72(2),



No of participants	48		
sample/unit	A	µmol/l	B
mean	138		116
standard deviation	31.7		29.5
coefficient of variation	22.9		25.3

Sample A [µmol/l]

M	Kit	N	Min	16.P	50.P	84.P	Max
Alle	48	63.0	105	141	167	282	
1	99	18	82.2	101	143	156	190
1	117	7	141		157	282	
2	99	7	63.0		122	193	
4	99	6	99.9		137	167	

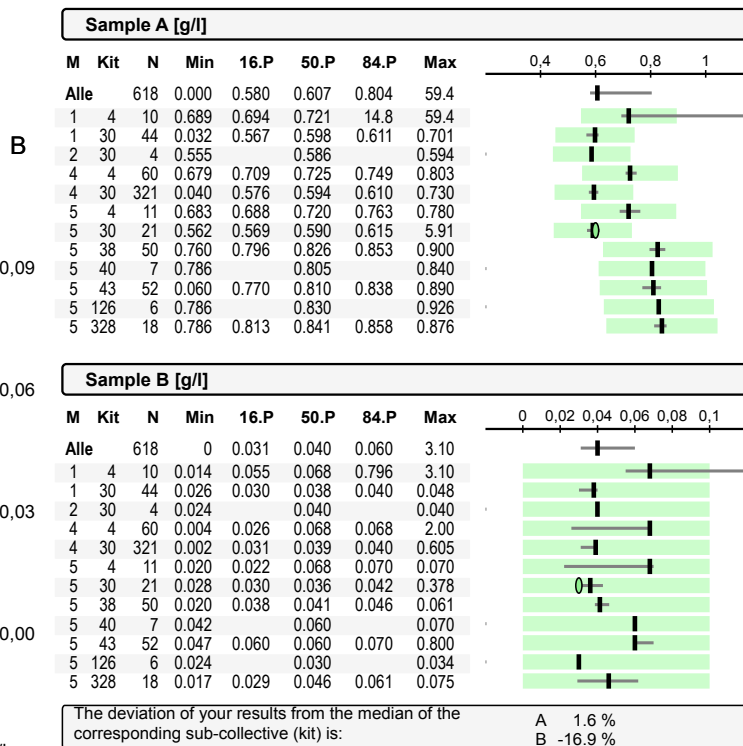
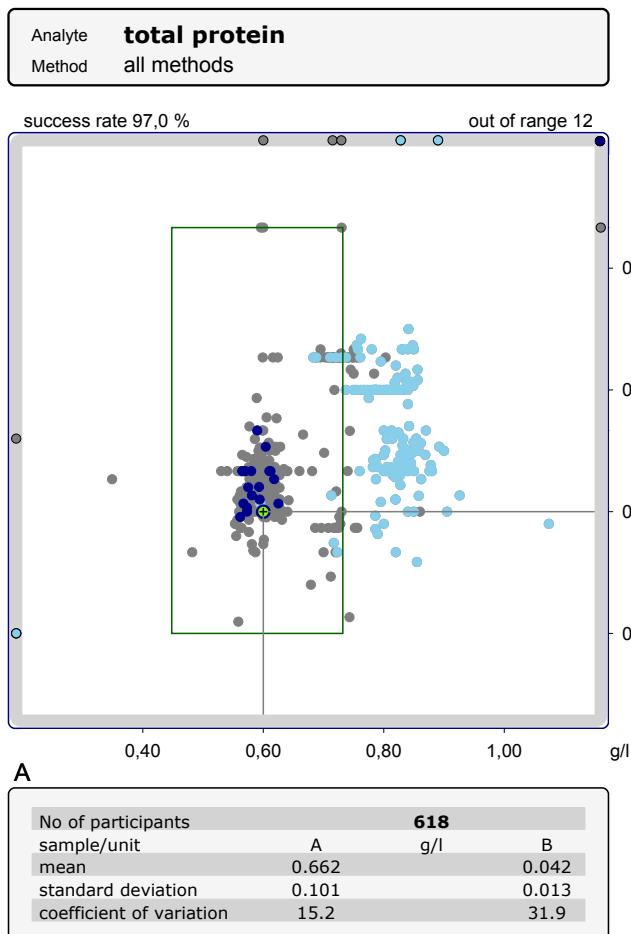
Sample B [µmol/l]

M	Kit	N	Min	16.P	50.P	84.P	Max
Alle	48	63.0	99.0	112	146	343	
1	99	18	73.3	100	113	130	187
1	117	7	84.4		139	343	
2	99	7	65.0		102	170	
4	99	6	76.0		112	122	

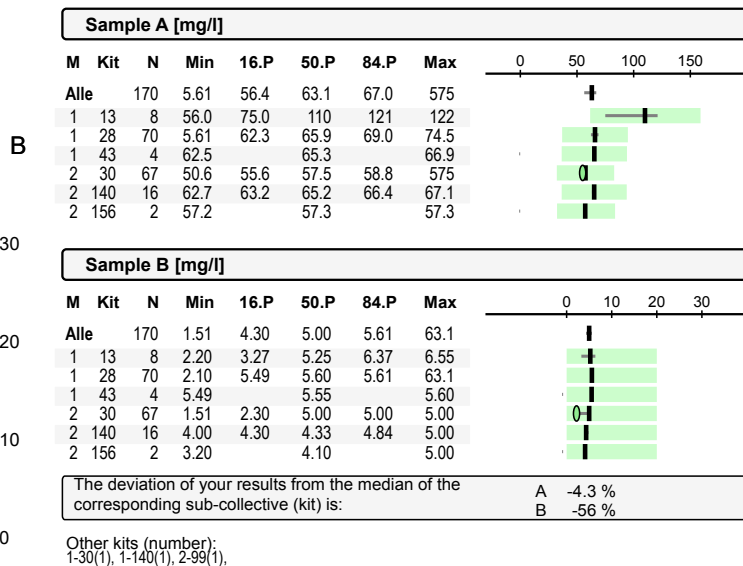
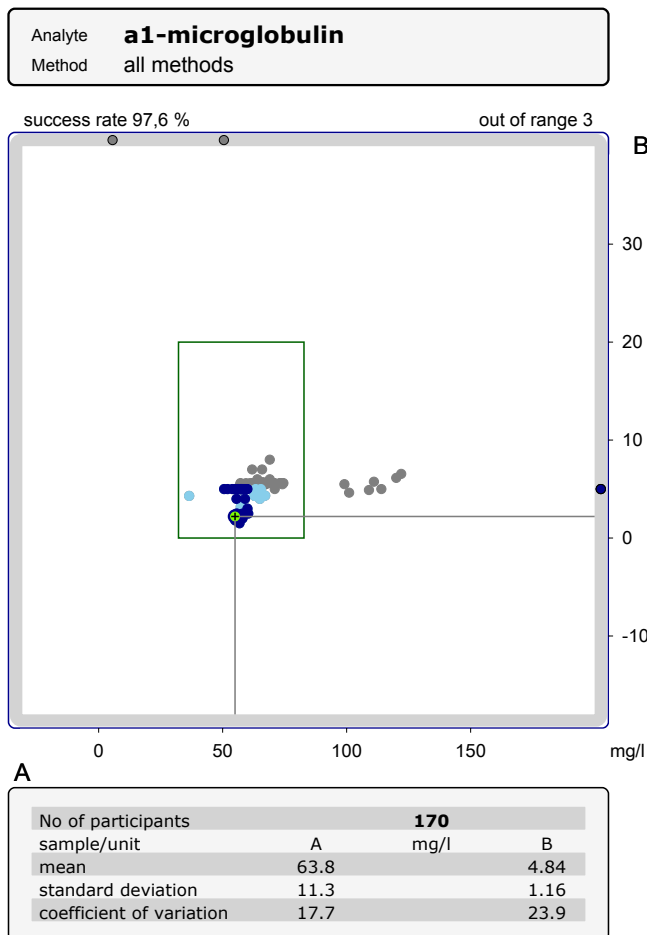
The deviation of your results from the median of the corresponding sub-collective (kit) is:

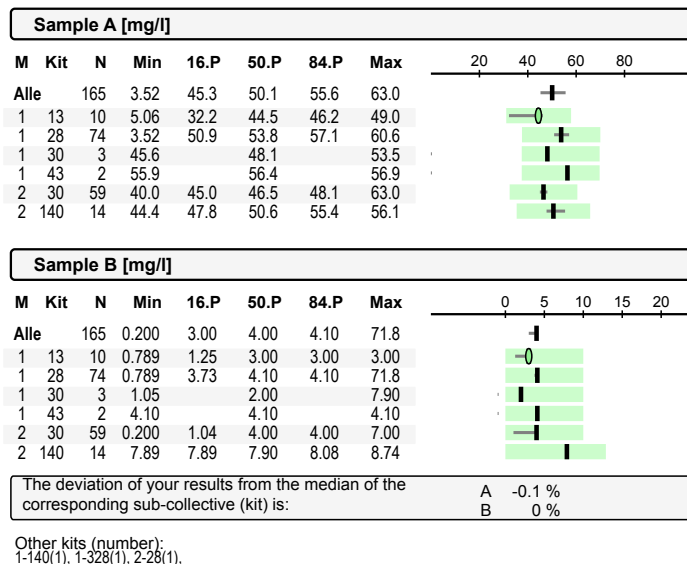
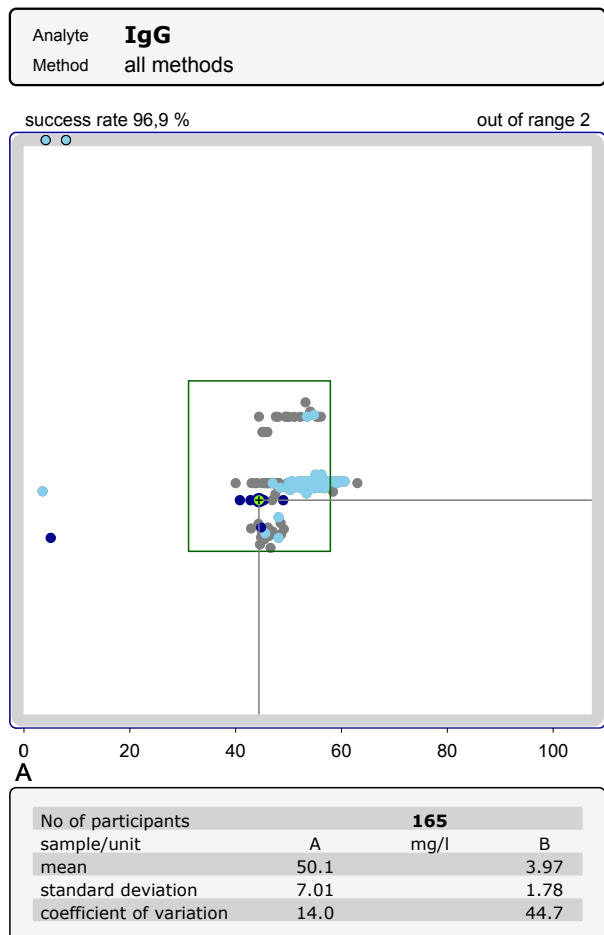
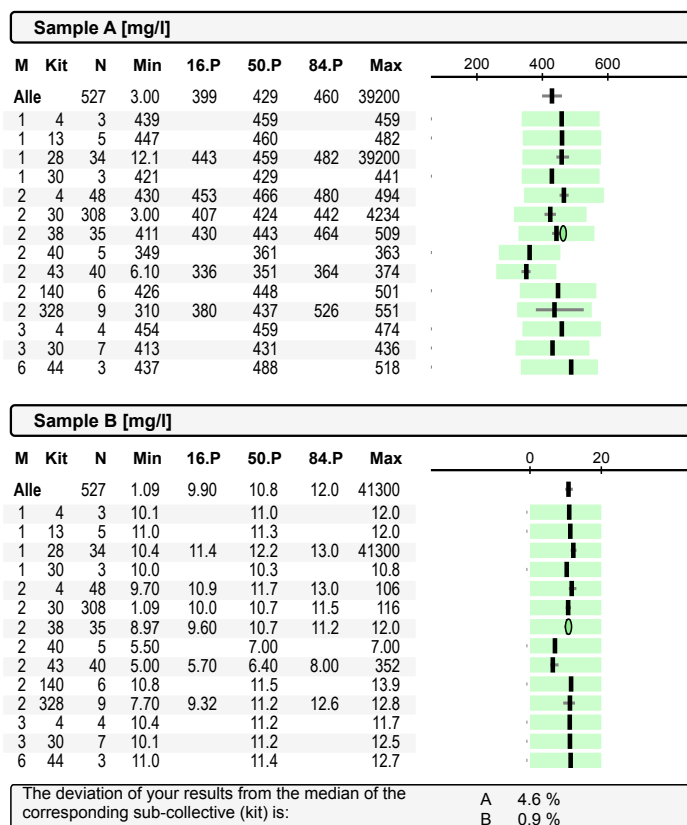
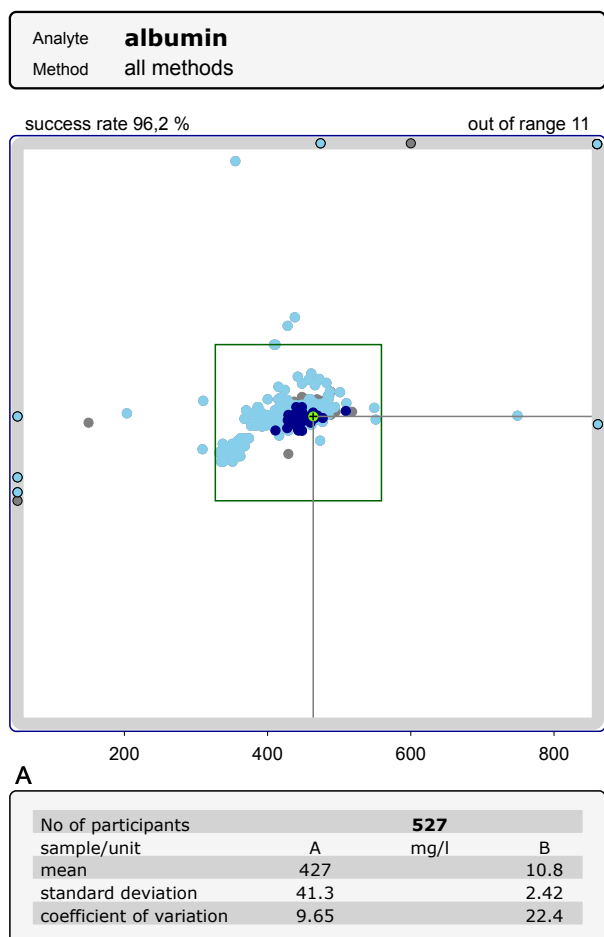
A 32 %
B 65 %

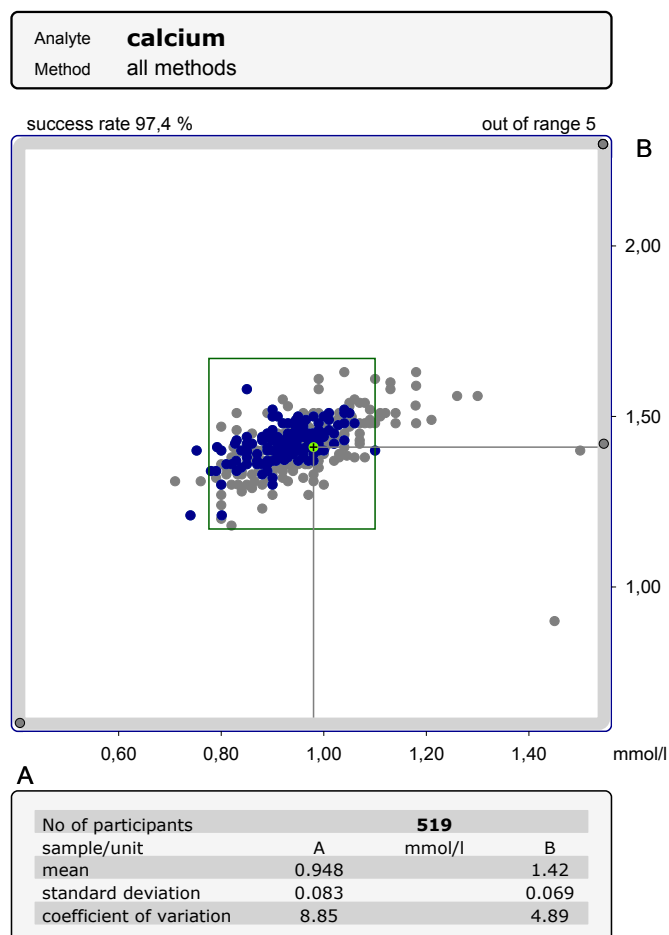
Other kits (number):
1-30(1), 1-38(2), 1-126(1), 1-168(1), 2-39(1), 4-12(1), 4-72(2), 4-168(1),



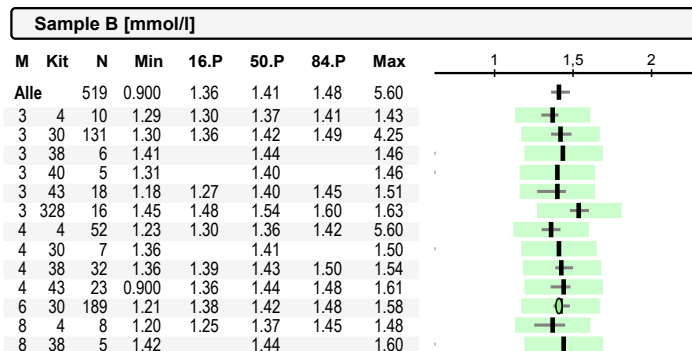
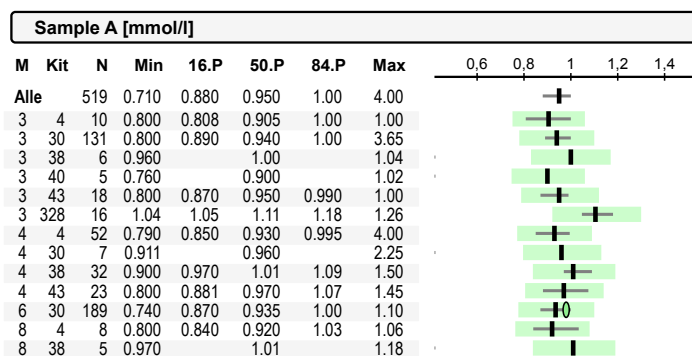
Other kits (number):
1-43(1), 1-126(1), 2-04(1), 3-30(2), 4-43(1), 5-13(1), 5-53(1), 5-105(1), 5-108(1), 5-128(2), 7-40(1), 8-08(1),







B

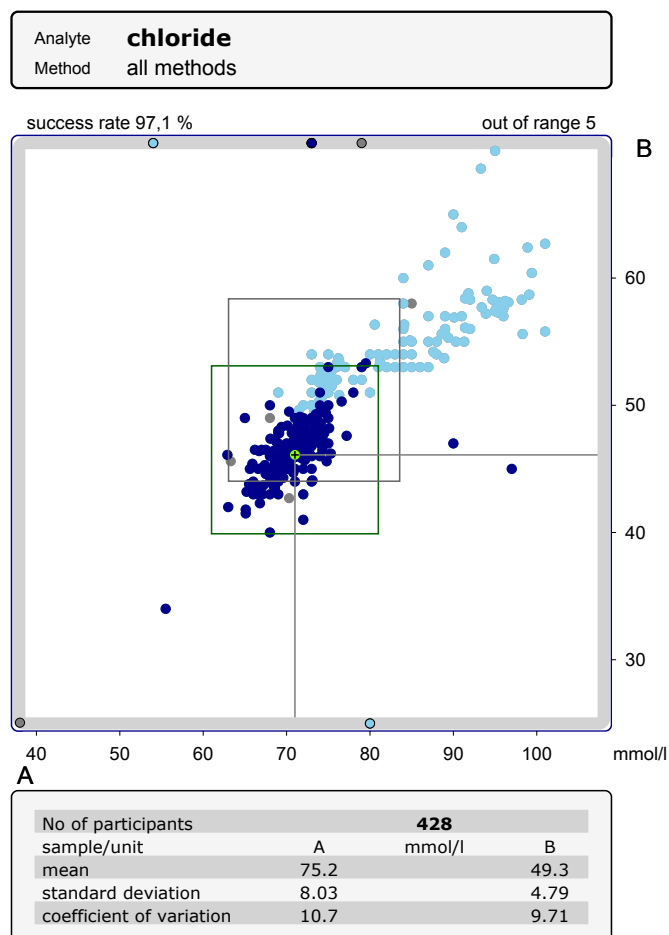


The deviation of your results from the median of the corresponding sub-collective (kit) is:

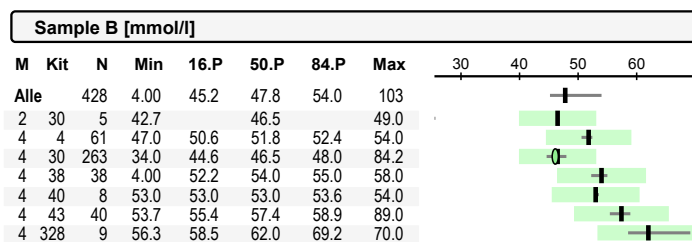
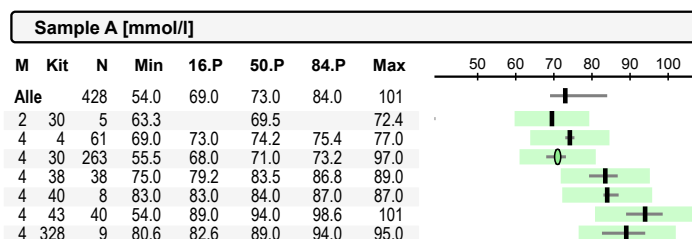
A 4.8 %
B -0.7 %

Other kits (number):

1-43(1), 1-91(1), 1-99(1), 1-134(1), 3-07(1), 3-128(1), 4-40(3), 4-126(1), 7-30(1), 7-38(1), 8-126(3), 8-134(2),



B



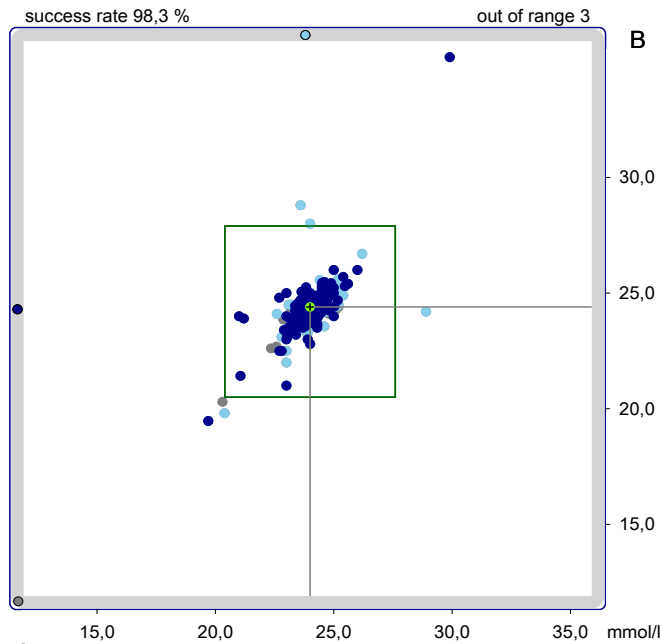
The deviation of your results from the median of the corresponding sub-collective (kit) is:

A 0 %
B -0.8 %

Other kits (number):

1-358(1), 2-328(1), 4-530(1), 4-128(1),

Analyte **potassium**
Method all methods



No of participants	601		
sample/unit	A	mmol/l	B
mean	24.1		24.3
standard deviation	0.740		0.833
coefficient of variation	3.07		3.43

Probe A (TV = 24.0 [mmol/l])

M	Kit	N	Min	16.P	50.P	84.P	Max		15	20	25	30
Alle	601		4.50	23.6	24.0	24.8	29.9					
2	4	69	22.6	23.5	23.9	24.1	25.2					
2	30	366	4.50	23.6	24.0	24.6	29.9					
2	38	47	23.0	24.0	24.0	25.0	28.9					
2	40	10	23.5	23.9	24.1	24.9	25.0					
2	43	45	23.2	24.4	25.0	25.2	26.2					
2	328	20	20.4	23.1	23.7	24.0	24.5					
3	30	8	23.4	23.5	24.1	24.8	24.9					
4	4	5	23.7		24.2		24.7					
4	30	16	22.9	23.1	23.9	24.6	25.0					
4	38	3	24.0		24.0		24.3					

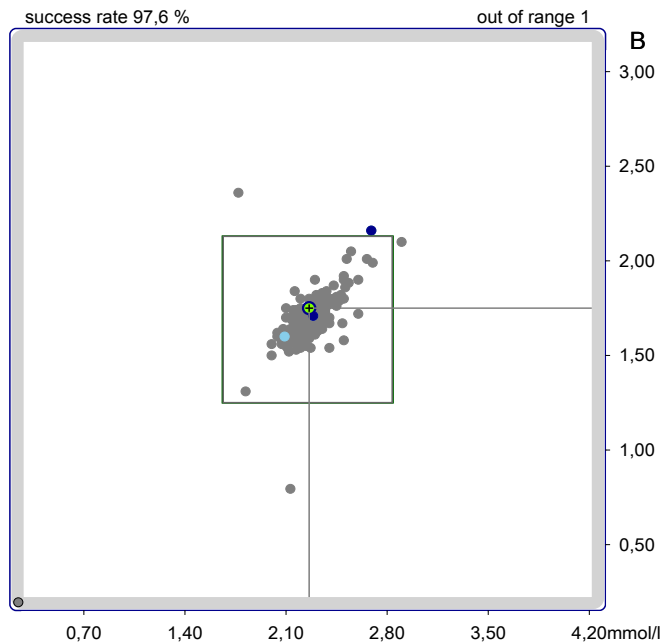
Probe B (TV = 24.2 [mmol/l])

M	Kit	N	Min	16.P	50.P	84.P	Max		15	20	25	30
Alle	601		19.5	23.9	24.2	25.0	62.0					
2	4	69	23.3	23.7	24.1	24.6	28.8					
2	30	366	19.5	23.9	24.2	24.9	35.2					
2	38	47	23.5	24.0	24.2	25.0	28.0					
2	40	10	23.6	23.8	24.3	24.9	25.1					
2	43	45	23.5	24.3	24.9	25.2	26.7					
2	328	20	19.8	22.8	23.8	24.4	62.0					
3	30	8	23.7	23.7	24.0	25.2	25.4					
4	4	5	24.0		24.5		24.8					
4	30	16	23.0	23.8	24.4	25.0	25.0					
4	38	3	24.0		24.0		24.5					

The deviation of your results from the median of the corresponding sub-collective (kit) is:

Other kits (number):
1-07(1), 2-128(1), 3-04(2), 3-07(1), 3-38(1), 3-43(1), 3-126(2), 4-43(2), 5-30(1),

Analyte **magnesium**
Method all methods



No of participants	254		
sample/unit	A	mmol/l	B
mean	2.26		1.69
standard deviation	0.133		0.123
coefficient of variation	5.91		7.24

Probe A (TV = 2.25 [mmol/l])

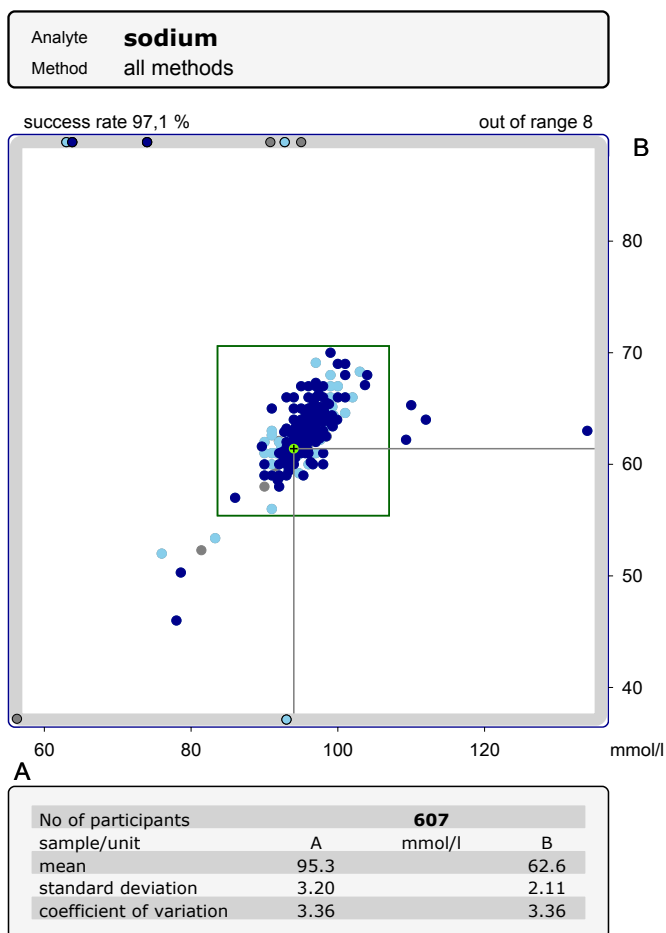
M	Kit	N	Min	16.P	50.P	84.P	Max		0,7	1,4	2,1	2,8	3,5
Alle	254		1.77	2.16	2.25	2.35	2.90						
1	99	3	2.26		2.29		2.69						
2	4	9	2.20	2.21	2.28	2.42	2.50						
2	30	116	1.82	2.13	2.24	2.38	2.90						
2	38	15	2.08	2.14	2.26	2.30	2.55						
2	40	6	2.09		2.29		2.35						
2	43	26	1.77	2.16	2.25	2.35	2.52						
3	4	25	2.07	2.17	2.27	2.31	2.47						
4	30	3	2.20		2.25		2.35						
5	30	19	2.15	2.19	2.27	2.37	2.90						
6	4	4	2.22		2.28		2.32						
6	30	8	2.14	2.14	2.22	2.28	2.30						
6	43	6	2.13		2.25		2.38						
6	328	4	2.16		2.27		2.38						

Probe B (TV = 1.69 [mmol/l])

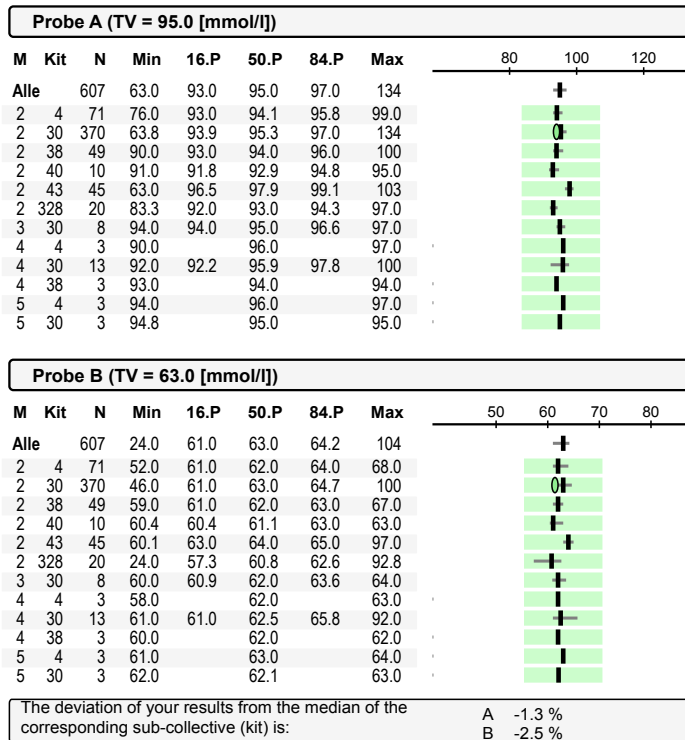
M	Kit	N	Min	16.P	50.P	84.P	Max		1	1,5	2	2,5
Alle	254		0.795	1.60	1.69	1.76	2.36					
1	99	3	1.71		1.75		2.16					
2	4	9	1.60	1.60	1.64	1.84	1.92					
2	30	116	1.31	1.60	1.68	1.79	2.10					
2	38	15	1.59	1.60	1.64	1.71	2.05					
2	40	6	0.795		1.70		1.83					
2	43	26	1.58	1.60	1.71	1.79	2.36					
3	4	25	1.56	1.64	1.70	1.76	1.80					
4	30	3	1.60		1.69		1.72					
5	30	19	1.61	1.64	1.70	1.78	2.10					
6	4	4	1.64		1.70		1.73					
6	30	8	1.58	1.59	1.63	1.70	1.70					
6	43	6	1.63		1.70		1.80					
6	328	4	1.63		1.66		1.68					

The deviation of your results from the median of the corresponding sub-collective (kit) is:

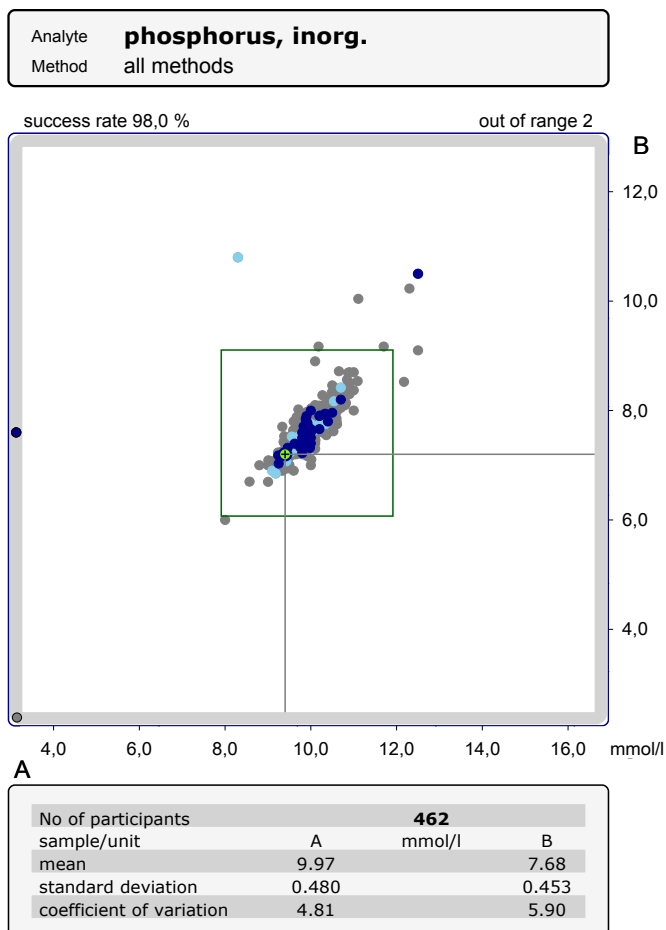
Other kits (number):
1-91(1), 2-328(1), 3-30(1), 4-38(1), 6-38(1), 6-91(1), 6-99(2), 6-134(2),



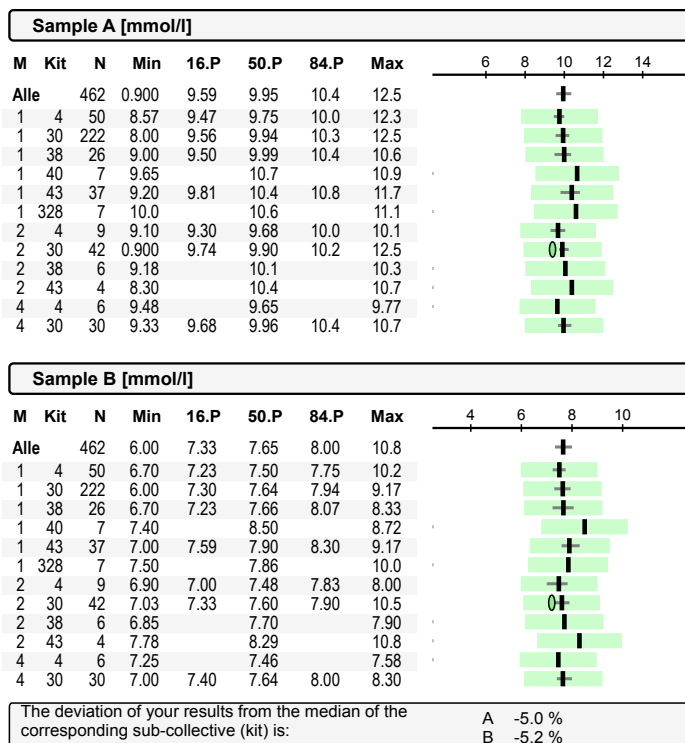
B



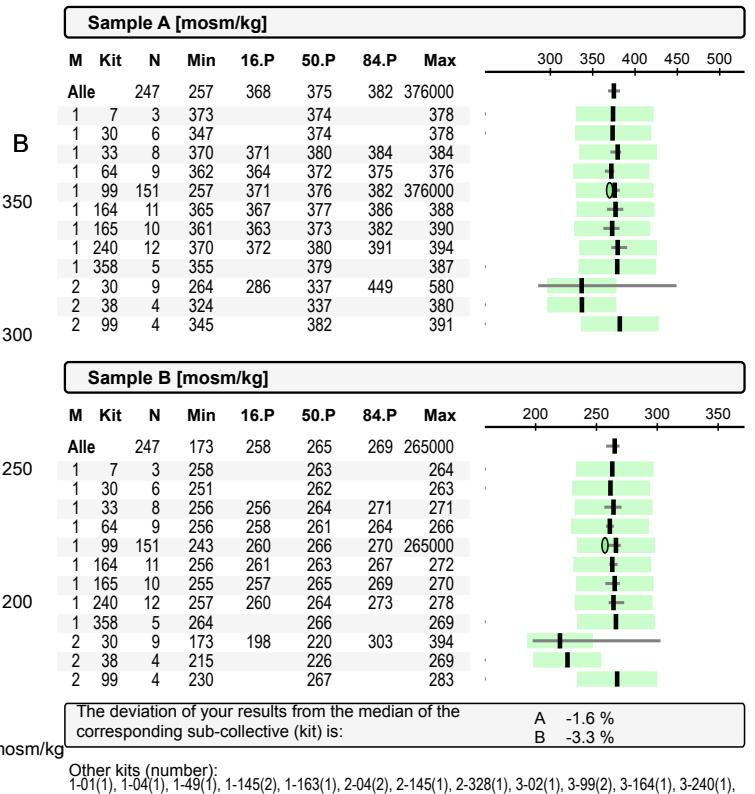
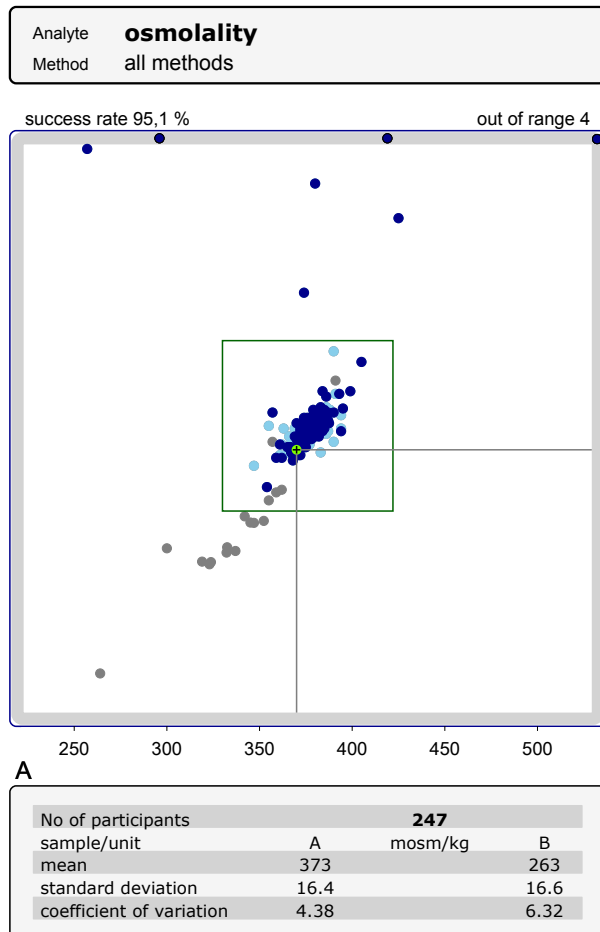
Other kits (number):
1-07(1), 1-30(1), 2-128(1), 3-04(2), 3-43(1), 3-126(1), 4-43(2),



B



Other kits (number):
1-07(1), 1-126(3), 1-128(1), 3-08(1), 3-38(1), 4-38(3), 4-40(1), 4-43(1), 4-99(1), 4-328(3),



Analyte **proteinuria**
Method all methods

- evaluation of diagnosis -

Sample A

- normal excretion of proteins
- glom. proteinuria (a1-microglobulin <14mg/g crea!)
- 124 glomerular and tubular proteinuria
- 1 tubular proteinuria (e.g. tubular interst. nephropathy)
- prerenal proteinuria (e.g. Bence-Jones proteinuria)
- postrenal proteinuria (e.g. in postrenal blood flow)
- other constellations

Sample B

- 124 normal excretion of proteins
- 1 glom. proteinuria (a1-microglobulin <14mg/g crea!)
- glomerular and tubular proteinuria
- tubular proteinuria (e.g. tubular interst. nephropathy)
- prerenal proteinuria (e.g. Bence-Jones proteinuria)
- postrenal proteinuria (e.g. in postrenal blood flow)
- other constellations

number of participants

125

[success rate 98 %]

Analyte **pH ST**
Method all methods

Sample A

-	-	pH value 5
-	1	pH value 5,5
1	2	pH value 6
3	8	pH value 6,5
15	157	pH value 7
22	273	pH value 7,5
23	210	pH value 8
-	-	pH value 8,5
-	-	pH value 9

Sample B

1	1	pH value 5
1	1	pH value 5,5
3	5	pH value 6
3	25	pH value 6,5
42	402	pH value 7
12	160	pH value 7,5
2	57	pH value 8
-	-	pH value 8,5
-	-	pH value 9

visuell optisch-automatisiert

number of participants

715

Analyte **protein ST**
Method all methods

sample A

sample A	2 30	2 145	2 40	2 146	1 30	others	all	method kit
1	1	1	2	-	-	0	4	negative
2	1	1	3	-	3	5	14	weak/borderline pos.
98	23	23	7	28	9	54	219	positive
131	128	128	114	4	25	63	465	strong positive

sample B

sample B	2	2	2	2	1	others	all	method
	30	145	40	146	30			kit
	217	150	126	32	34	120	679	negative
	8	1	-	-	3	0	12	weak/borderline pos.
	6	2	-	-	-	1	9	positive
	1	-	-	-	-	0	1	strong positive

number of participants

702

Analyte **glucose ST**
Method all methods

sample A

sample A	2 30	2 145	2 40	2 146	1 30	others	all	method kit
1	-	2	-	-	0	3	< 50 mg/dl	
17	28	34	17	1	37	134	50-99 mg/dl	
211	122	89	15	21	80	538	100-299 mg/dl	
2	-	1	-	15	6	24	300- 999 mg/dl	
-	2	-	-	1	0	3	>=1000 mg/dl	

sample B

sample B	2 30	2 145	2 40	2 146	1 30	others	all	method kit
	230	152	126	32	38	122	700	< 50 mg/dl
	1	-	-	-	-	0	1	50-99 mg/dl
	-	-	-	-	-	0	0	100-299 mg/dl
	-	-	-	-	-	0	0	300- 999 mg/dl
	-	-	-	-	-	0	0	>=1000 mg/dl

number of participants

702

Analyte **hemoglobin ST**
Method all methods

sample A

sample A	2	2	2	2	1	others	all	method kit
	30	145	40	146	30			
</								

sample B

sample B	2 30	2 145	2 40	2 146	1 30	others	all	method kit
	227	152	125	31	40	122	697	negative
	1	1	-	1	-	0	3	5 - 12 Erys/ul
	3	-	-	-	2	0	5	13 - 50 Erys/ul
	-	-	-	-	-	1	1	> 50 Erys/ul

number of participants

706

Analyte leukocytes ST		Method all methods						
sample	2	2	2	2	1	others	all	method kit
A	30	145	40	146	30			
	230	152	123	32	39	123	699	negative
	1	1	-	-	2	1	5	5 - 50 /ul
	-	-	-	-	-	0	0	51 - 250 /ul
	-	-	-	-	-	0	0	> 250 /ul

Analyte keton ST		Method all methods						
sample	2	2	2	2	1	others	all	method kit
A	30	145	40	146	30			
	232	153	124	31	36	121	697	negative/trace
	-	-	-	-	-	0	0	5 - 49 mg/dl
	-	-	-	-	-	0	0	>=50 mg/dl

sample	2	2	2	2	1	others	all	method kit
B	30	145	40	146	30			
	216	147	114	32	38	118	665	negative
	15	6	9	-	3	7	40	5 - 50 /ul
	-	-	-	-	-	0	0	51 - 250 /ul
	-	-	-	-	-	0	0	> 250 /ul

sample	2	2	2	2	1	others	all	method kit
B	30	145	40	146	30			
	231	153	124	31	36	121	696	negative/trace
	1	-	-	-	-	0	1	5 - 49 mg/dl
	-	-	-	-	-	0	0	>=50 mg/dl

number of participants

705

number of participants

697

Analyte nitrite ST		Method all methods						
sample	2	2	2	2	1	others	all	method kit
A	30	145	40	146	30			
	231	153	124	32	38	121	699	negative
	1	-	-	-	-	0	1	positive

Analyte bilirubin ST		Method all methods						
sample	2	2	2	2	1	others	all	method kit
A	30	145	40	146	30			
	230	153	121	32	37	120	693	negative / light
	-	-	-	-	-	0	0	middel
	-	-	-	-	-	0	0	strong

sample	2	2	2	2	1	others	all	method kit
B	30	145	40	146	30			
	232	152	124	32	38	121	699	negative
	-	1	-	-	-	0	1	positive

sample	2	2	2	2	1	others	all	method kit
B	30	145	40	146	30			
	230	153	121	32	37	120	693	negative / light
	-	-	-	-	-	0	0	middel
	-	-	-	-	-	0	0	strong

number of participants

700

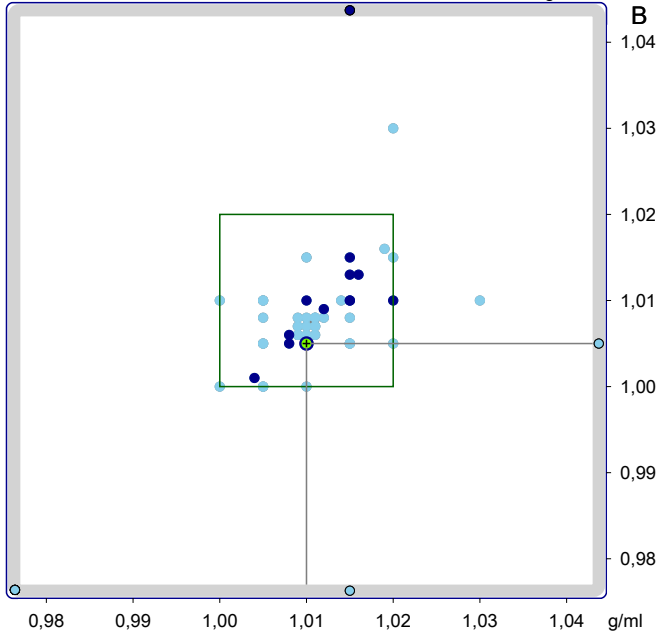
number of participants

693

Analyte **specific grav. ST**
Method all methods

success rate 97,7 %

out of range 13

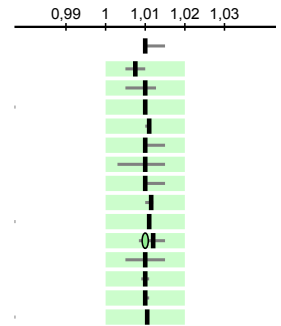


A

No of participants	675		
sample/unit	A	g/ml	B
mean	1.01		1.01
standard deviation	0.003		0.003
coefficient of variation	0.315		0.299

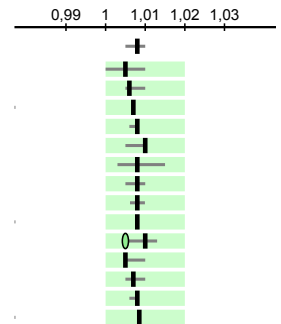
Sample A [g/ml]

M	Kit	N	Min	16.P	50.P	84.P	Max
Alle	675	0.001	1.01	1.01	1.02	1.10	
1	30	22	0.001	1.01	1.01	1.01	1.02
1	99	10	1.01	1.01	1.01	1.01	1.02
1	145	7	1.01		1.01		1.01
2	13	16	1.01	1.01	1.01	1.01	1.01
2	30	225	0.001	1.01	1.01	1.02	1.10
2	33	9	1.00	1.00	1.01	1.02	1.02
2	40	123	0.001	1.01	1.01	1.02	1.03
2	43	12	1.01	1.01	1.01	1.01	1.01
2	64	7	1.01		1.01		1.02
2	99	19	1.00	1.01	1.01	1.02	1.02
2	127	10	1.01	1.01	1.01	1.02	1.02
2	145	149	0.001	1.01	1.01	1.01	1.02
2	146	33	1.01	1.01	1.01	1.01	1.02
2	164	6	1.01		1.01		1.02



Sample B [g/ml]

M	Kit	N	Min	16.P	50.P	84.P	Max
Alle	675	0.001	1.01	1.01	1.01	1.10	
1	30	22	0.001	1.00	1.01	1.01	1.01
1	99	10	1.01	1.01	1.01	1.01	1.01
1	145	7	1.01		1.01		1.01
2	13	16	1.01	1.01	1.01	1.01	1.01
2	30	225	0.001	1.01	1.01	1.01	1.02
2	33	9	1.00	1.00	1.01	1.02	1.02
2	40	123	0.001	1.01	1.01	1.01	1.03
2	43	12	1.01	1.01	1.01	1.01	1.01
2	64	7	1.01		1.01		1.02
2	99	19	1.00	1.01	1.01	1.01	1.10
2	127	10	1.01	1.01	1.01	1.01	1.01
2	145	149	0.001	1.01	1.01	1.01	1.02
2	146	33	1.01	1.01	1.01	1.01	1.01
2	164	6	1.01		1.01		1.02



The deviation of your results from the median of the corresponding sub-collective (kit) is:

A -0.2 %
B -0.5 %

Other kits (number):
1-12(1), 1-34(1), 1-40(5), 1-64(1), 1-127(2), 1-151(1), 2-02(1), 2-12(1), 2-17(1), 2-25(1), 2-28(2), 2-34(3), 2-37(5), 2-140(1), 2-211(1),

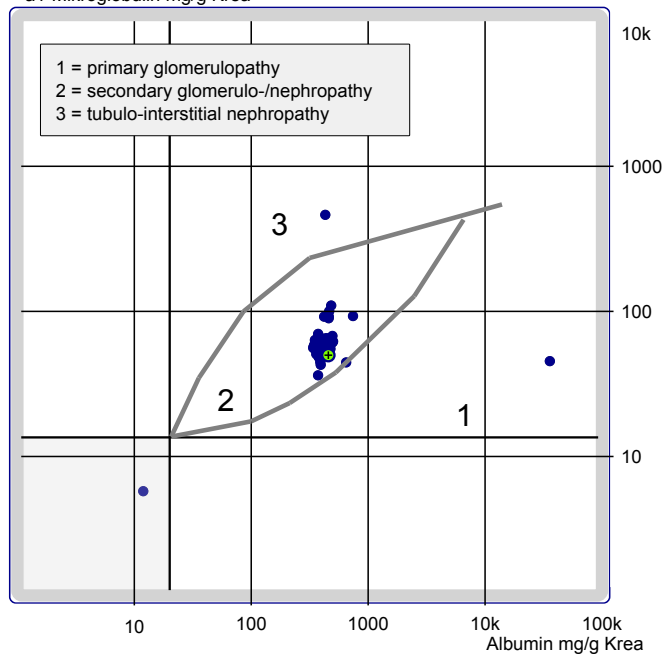
Analyte **urobilinog. ST**
Method all methods

sample	2	2	2	2	1	others	all	method
A	30	145	40	146	30			kit

sample	2	2	2	2	1	others	all	method
B	30	145	40	146	30			kit

Analyte **diagn. graph A**
Method

a1-Mikroglobulin mg/g Krea

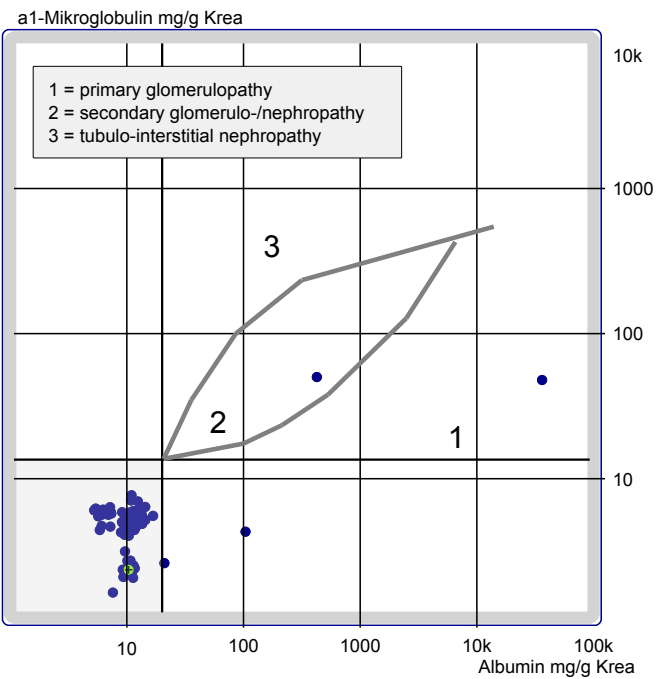


No of participants	152		
abscissa/ordinate	Albumin	mg/g Krea	a1-Mikrogl.
median	429		61.4

number of participants

693

Analyte **diagn. graph B**
Method



No of participants		152	
abscissa/ordinate	Albumin	mg/g Krea	a1-Mikrogl.
median	10.4		4.88